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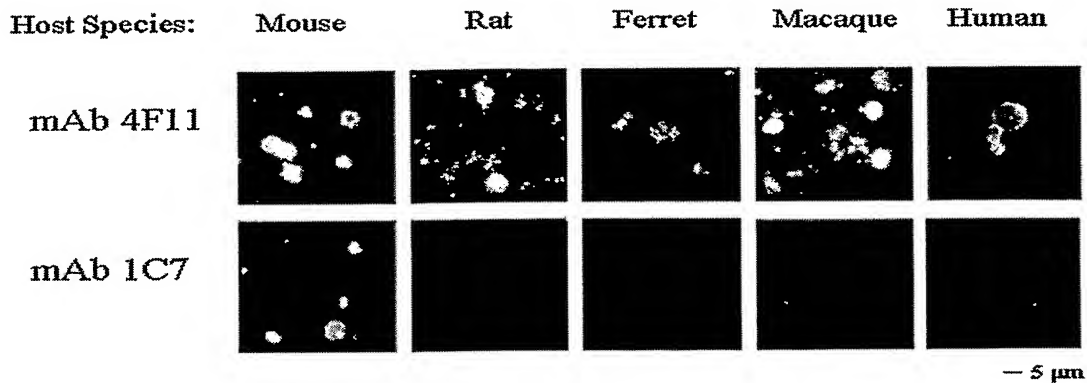


Figure 1

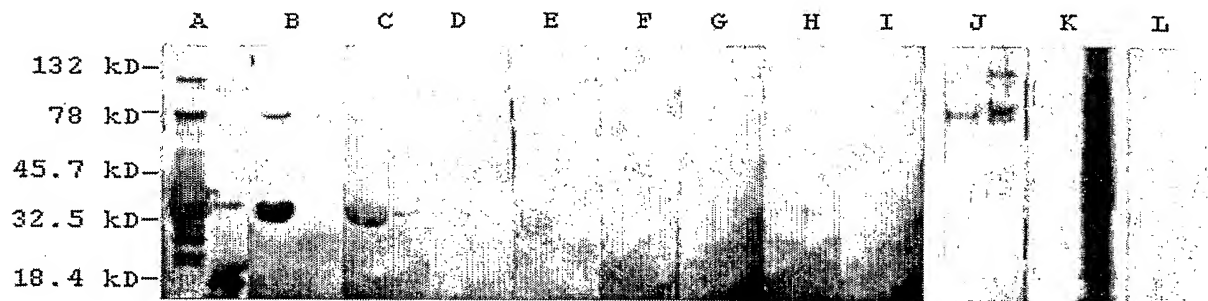
K P T P Q P T P Q P T S E P T S E P T S E P T S E P T P Q P A P P Q 735
2113 AAACCAACACCTCAACCAACCTCAGCCAACATCTGAGCCAACATCTGAGCCAACATCTGAACCAACACCTCAACCAGCACCTC
P A P P Q P A P Q P A P Q P A P Q P A P P Q P A P P Q P V P P Q P 768
2213 AACCAGCACCTCAACCAGCACCTCAACCAGCACCTCAACCAGCACCTCAACCAGCACCTCAACCAGTACCTCAACC
V P P Q P M P S R P A P P K P T P Q P T S E P A P Q P T S E S T S 801
2313 AGTACCACCTCAACCAATGCCATCTAGACCAGCACCTAAACCAACACCTCAACCAACATCTGAGCCAGCACCTCAACCAACATCTGAGTCAACATCT
E P T P R P P P Q P T S E P T S E P T S E P T S E P S P Q P T P Q P 835
2413 GAACCAACACCTCGACCACCCTCAGCCAACATCTGAGCCAACATCTGAACCAACATCTGAACCAACATCTGAACCAACCTCAACCAACACCTCAAC
V P Q P A P Q P A P P K P A P K P T P P K P A P K P T P P K P A P 868
2513 CAGTACCTCAACCAGCACCTCAACCAGCACCTAAACCGGCACCTAAACCAACACCACCTAAACCGGCACCTAAACCAACACCACCTAAACCGGC
K P A P S K S S S K P T S T
2613 TAAACAGCACCATCTAAATCATCATCTAAACCAACATCTACA

Figure 2A

T N I S E P A L P D K D P Q P T S S P Q P K P R P R P R P Q P Q P H 34
1 ACCAATATATCCGAACAGCACTGCCTGATAAGGATCCTCAACCTACATCTTCACCTCAGCCAAAACCTCGGCCAAGACCTCGACCTCAACCTCAACCTC
P H P K P Q P Q P T P E P Q P Q P A P E P R P Q P T S K P R P Q P 67
101 ATCCACATCCAAAACCTCAGCCTCAGCCGACGCCAGAACCTCAGCCTCAGCCGGCGCCAGAACCTCGACCTCAGCCGACGTCAAAAACCTCAGCCTCAGCC
T S K P R P Q P T P E P R P L P V P G P G P L P V P G P R P Q P Q 100
201 AACGTCAAAACCTCGACCTCAGCCGACGCCAGAACCTCGACCTCTGCCGGTGCCAGGACCTGGACCTCTGCCGGTGCCAGGACCTCGACCTCAACCTCAA
P Q P Q P Q P Q P Q P Q P Q P Q P Q P Q P Q P Q P Q P Q P K P Q 134
301 CCTCAACCTCAACCTCAGCCTCAACCTCAACCTCAGCCTCAACCTCAGCCTCAGCCTCAGCCTCAGCCTCAACCTCAGCCGAAGCCTC
P P S Q S T S E S A S Q S K P K P T T Q T K P S P R P H P K P V P 167
401 AAACCATCTCAGTCAACATCAGATCGCAATCCAACCAAAACCAACAACACAAACAAACCGTCACCGAGACCACACCAAAAGCCGGTGCC
K P S S I D T G P S K S D S S F I F T V T K T I T K I S E T E K P 200
501 AAAACCATCATCGATAGACACAGGACCATCAAAATCGGATTCAGGCTCATTTTACAGTAACAAAACAATAACAAAGATATCAGAAACAGAAAACCA
S T K P S V K P T S T K T T S K P S T K P S T K P S V K P A S T K T 234
601 TCTACAAAACCATCTGTGAAACCAACCTCTACAAAGACAACATCAAAACCATCTACAAAACCATCTGTAAACACGCTCTACAAAGA
T S E S E K P T L E E V P E T K G N G V R V I G F E G L Q L L S M 267
701 CAACATCAGAAATCAGAAAACCAACATTGGAAGAAGTTCCAGAAACTAAAGGGAATGGTGTAAGAGTAATAGGATTTGAGGGGTACAAATATTATCAAT
I V A I I I G I W I M *
801 GATTGTTGCAATAAATTGGGATATGGATAATGTAATTTAATTAGAAGTCATTGGCTATTAAATTAATATATAGTAATTTGTAATAATTAGATAATA
901 GACAGGGGATCTAGAAATCAATGTGTGATTAAATAAATATAAAAAATCAAAAAAAAAAAAAAAAAA

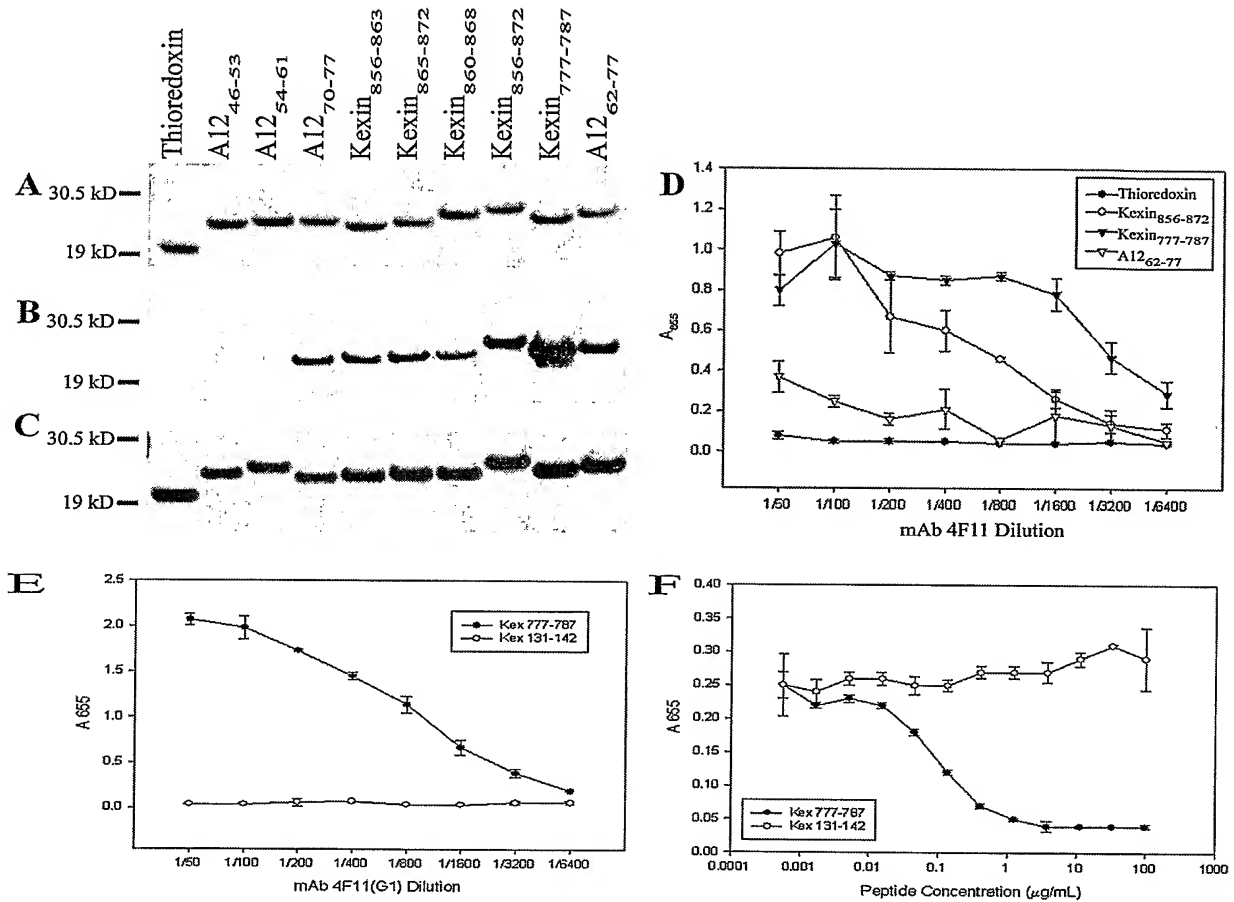
Figure 2B

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**Figure 3**

Construct	Sequence of insert
KEXIN ₈₅₆₋₈₇₂	KPAPKPTPPKPAPKPAP
KEXIN ₈₅₆₋₈₆₃	KPAPKPTP
KEXIN ₈₆₅₋₈₇₂	KPAPKPAP
KEXIN ₈₆₀₋₈₆₈	KPTPPKPAP
KEXIN ₇₇₇₋₇₈₇	RPAPPKPTPQP
A12 ₄₆₋₅₃	EPQPQPAP
A12 ₅₄₋₆₁	EPRPQPTS
A12 ₇₀₋₇₇	KPRPQPTP
A12 ₆₂₋₇₇	KPRPQPTS-KPRPQPTP

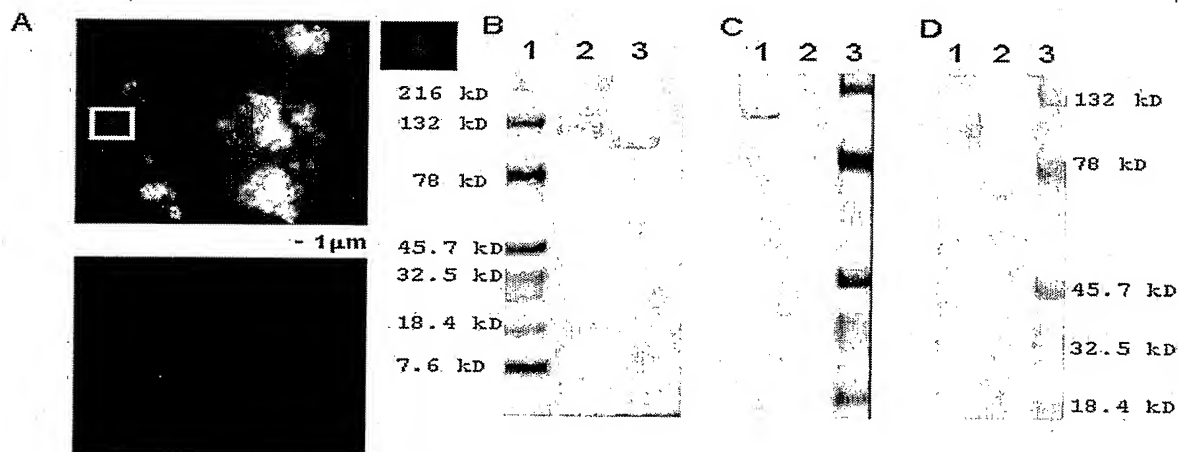
Figure 4



Figures 5A-F

URSP2 PspA	<u>EKELKEIDESDSEDIKEGLRAPLOSKLDAKKAKLSKLEELSDKIDEL</u>	280
URSP2 PspA	<u>DAEIAKLEKDVEDFKNSDGEQAEQYLVAAKKDLDAKKAELENTEADLK</u>	328
Kex ₈₅₆₋₈₇₂		
	KPAPKPTP.PKPAPKPAP	
URSP2 PspA	<u>KAVDEPETPAPAPKPAPAPAPTPEAPAPAPKPAPAPKPAPAPAPTPEA</u>	376
	:	
Kex ₇₇₇₋₇₈₇		
	RPAP.PKPTPQP	
Kex ₈₅₆₋₈₇₂	KPAPKPTP.PKPAPKPAP	
URSP2 PspA	<u>PAPAPKPAPAPKPAPAPAPTPEAPAPAPKPAPAPRPAPAPKPAPDPKP</u>	424
	:	
Kex ₇₇₇₋₇₈₇	RPAP.PKPTPQP	
	RPAP.PKPTPQP	

Figure 6



Figures 7A-D

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1 21 41 61 81
| | | | |
CTAGATACTCGTGCTAATGTATTTTCTTCATGTTATAAAGAAGATATGGATTTTTCAGCCAAATTAGATCTTCTAAATAGGATAAAAGAT
L D T R A N V F S S C Y K E D M D F S A K L D L L N R I K D

101 121 141 161
| | | |
AAGATTGTAGTTCCAAAAGGAAACACGAGGTATTTTGTAGAGTTATTTGTGTAAAAGCTATATTGTCGCCGAATGCAGCGCCAGTGATTTA
K I V V P K G N T R Y F V E L L C K S Y I V A E C S A S D L

181 201 221 241 261
| | | | |
ATGTTCAAATCTTATGCTCTTATGGAAGCCTGTCTTCACCCAGAAAGGATCTGTAGAGAATTAAAAAATCATTTTTCCGAAGAATCTAGG
M F K S Y A L M E A C L H P E R I C R E L K N H F S E E S R

281 301 321 341
| | | |
AAATTAGAAAATAAATTAAGGAGTATTTTAAAACCCACATATTATGAATGCAAAGATCTAGGACAAAAGTGCAACTCTGGATTTTATTTT
K L E N K L R S I L K P T Y Y E C K D L G Q K C N S G F Y F

361 381 401 421 441
| | | | |
GATGGAGATATAGAAGCTCAATGCAATCATTTCAAAAAAAGATGTCAAGATAAACAAGAGAGACTAAAATTAATTAATCATATTGTTGAT
D G D I E A Q C N H F K K R C Q D K Q E R L K L I N H I V D

461 481 501 521
| | | |
TCATCTGCTCTTTATCTCGCAAATGAAGTACAATGCAGAACTTATTTTCGACAGTTTTTGTGGTGCGAATGTAAAACAAGAATTCAAACAA
S S A L Y L A N E V Q C R T Y F D S F C G A N V K Q E F K Q

541 561 581 601 621
| | | | |
ATATGCAACAAAGGAGCTAATGGCATATGCCCTGATATAATAGATGATTCTAAAGAACATTGTGCTCATTGATTAATCATTTAAACATCT
I C N K G A N G I C P D I I D D S K E H C A H L I N H L T S

641 661 681 701
| | | |
CTTGGAAATTTTCATGCTCTTCTGCTCACTTCCATTGGACTATTGCGACTCAGCGATTAATTACTGTAATTCFCTTTTCGAAGTTTTGCACG
L G I S S S S A S L P L D Y C D S A I N Y C N S L S K F C T

721 741 761 781 801
| | | | |
GAATCAAAACGACAGTGCATTCTGTTATTTCTTTCTGCACTAGCGAATCAAAAAAAGTGAATGAATATGGTTCTTTTATTGACCAATAT
E S K R Q C D S V I S F C T S E S K K T D E Y G S F I D Q Y

821 841 861 881
| | | |
CCCGCGGCTGCAGCAAATGCAACCAAATGCAAGGTAACCTTGAAGAGTTATGCCAAGATTCAAGCAAAAAGACTCTTATTCAACACTA
P A A A A N A T K C K V T L K E L C Q D S S K K D S Y S T L

901 921 941 961 981
| | | | |
TGTGCTTATAATAAAGATGGTTATACCGAAATATGTAAAAACTTAAGAAATTCATAGAAAAAGCATGCGAGAATTTGAGAATTCATTTA
C A Y N K D G Y T E I C K N L R N F I E K A C E N L R I H L

1001 1021 1041 1061
| | | |
CATACTTATGATACAAACTCACTCAATACGAATAAAGGATCTGCTCAAGATAGATGCACTTATATAAGAAATCTTTACTTTAAATTTAAA
H T Y D T N S L N T N K G S A Q D R C T Y I R N L Y F K F K

1081 1101 1121 1141 1161
| | | | |
AATATATGTTTATTGGTTGATCCTTTCTATGACTTATCTCCTATTATCACTCAAGAATGTAAAACCAATATATCCGAACCGACTGCCT
N I C L L V D P F Y D L S P I I T Q E C K T N I S E P A L P

1181 1201 1221 1241
| | | |
GATAAGGATCCTCAACCTACATCTTCACCTCAGCCAAAACCTCGGCCAAGACCTCGACCTCAACCTCAACCTCATCCACATCCAAAACCT
D K D P Q P T S S P Q P K P R P R P R P Q P Q P H P H P K P

Figure 8A

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1261 1281 1301 1321 1341
| | | | |
CAGCCTCAGCCGACGCCAGAACCTCAGCCTCAGCCGGCGCCAGAACCTCAGCCTCAGCCGACGTCAAAACCTCGACCTCAGCCAACGTCA
Q P Q P T P E P Q P Q P A P E P R P Q P T S K P R P Q P T S

1361 1381 1401 1421
| | | |
AAACCTCAGCCTCAGCCGACGCCAGAACCTCAGCCTCTGCCGGTGCCAGGACCTGGACCTCTGCCGGTGCCAGGACCTCAGCCTCAACCT
K P R P Q P T P E P R P L P V P G P G P L P V P G P R P Q P

1441 1461 1481 1501 1521
| | | | |
CAACCTCAACCTCAACCTCAGCCTCAACCTCAACCTCAGCCTCAACCTCAACCTCAGCCTCAGCCTCAGCCTCAGCCTCAACCT
Q P Q P Q P Q P Q P Q P Q P Q P Q P Q P Q P Q P Q P Q P

1541 1561 1581 1601
| | | |
CAGCCGAAGCCTCAACCACCATCTCAGTCAACATCAGAATCAGCATCGCAATCCAACCAAAACCAACAACACAAACAAACCGTCACCG
Q P K P Q P P S Q S T S E S A S Q S K P K P T T Q T K P S P

1621 1641 1661 1681 1701
| | | | |
AGACCACACCCAAAGCCGGTGCCAAAACCATCATCGATAGACACAGGACCATCAAAATCGGATTCAAGCTTCATTTTACAGTAACAAA
R P H P K P V P K P S S I D T G P S K S D S S F I F T V T K

1721 1741 1761 1781
| | | |
ACAATAACAAAGATATCAGAAACAGAAAAACCATCTACAAAACCATCTGTGAAACCAACCTCTACAAAGACAACATCAAAACCATCTACA
T I T K I S E T E K P S T K P S V K P T S T K T T S K P S T

1801 1821 1841 1861 1881
| | | | |
AAACCATCTACAAAACCATCTGTAAAACAGCCTCTACAAAGACAACATCAGAATCAGAAAAACCAACATTGGAAGAAGTTCAGAAACT
K P S T K P S V K P A S T K T T S E S E K P T L E E V P E T

1901 1921 1941 1961
| | | |
AAAGGAATGGTGAAGAGTAATAGGATTTGAGGGGTTACAATTATTATCAATGATTGTTGCAATAATAATTGGGATATGGATAATGTAA
K G N G V R V I G F E G L Q L L S M I V A I I I G I W I M -

Figure 8B